

Supplementary Table 1. *Arabidopsis* and rice protein sequences used for motif discovery with the MEME Suite.

Arabidopsis genes	Rice orthologs/sequelogs	Function	Protein sequences
At5g45350	No match	unknown function	MGGONDNDKDKGFHGYPAGYPPPGAYPPGPOQGYPPPGAYPPGAPGGYPPAGYGGYPPAGYGGYPPAGYGGYPPAGYPAHSGHAGGIGGMIAGAAAAYGAHHVAHSSSHGPYGAAYGHGFHGHYGHGKFKHGKFKHGKHMFGGGFKKKW
At1g47400	No match	unknown function	MMFSVANLAIKRFDHASTVVVEDVVDSSRVAYSENGDDDSGYD'APAA
At2g30760	No match	unknown function	MQMKPPVQRSHSHRRHLLRSLGHQFSGFFALCEKQRECKCDMATVMTTNLYNVARHSPHIIFVCGSWSSLIESAKEEEESMRRS
At2g30766	No match	unknown function	MAVVSHTNNAEGRLYESTQTWPIALYQIGGQENGDDDDCDVAPAA
At2g14247	No match	unknown function	MAPSAAMILSHPLVSHAKAKNQSLSSPSSVKSTRVFGFLWPWKALDNEDHSAVVLGRLFGDPATIEKRFQEALEQSCW
At1g13609	No match	unknown function	MNNLRVIMSVLLAVLVFTATVSESAEEMKGGEVTISLRCKTKECLKNIAEACVDCRCDKGICKCHGFTAETNNPTV
Rice locus	Arabidopsis ortholog/sequelogs	Function/description	Protein sequences
LOC_Os01g45914	#N/A	OslMA1	MAPVSEASPLVHDGGIiasFAVYAGAPCCSARGRMAETDGDDDDDYDCAPAA
LOC_Os03g51530	#N/A	Unknown function	MAPTASMLFLSYHQLHRPASAAEAPASRKEAEEEGSNGGAAGGRVRVSLLSALSLLARRREAAPTPTPQAAAKEVTRRGGGGDGVAGEGEPEEAASLERRFEAVRLSCWSS
LOC_Os11g15624	#N/A	Unknown function	METPSPPAHPCPOTSPNCRQGHWPNPSPALFSATTAGHQLRRSSITLKIPPDSQELKWTKKAMLKLKPATFHAEISFLRRRKKGKNSMVLTSSCYKSICSQLPVVFLSYFVGSEFVCRECWVPVGP
LOC_Os12g18410	#N/A	Unknown function	MPVTQHLSSEGDAQVQEATLAYSGAVAGVTIEELAGRRTSSWSVSYVLDLGFVHPHAAHACLLWERRAPCLPCPLQNIATAARPHLPPHTRLGLYISLTASLTTSCISCMGCVPCCHSTKEVSATAFEDNEAHSSRPEQGKVGEGLHVPGSEDQGREA
LOC_Os07g04920 - Os.12430.1.S1_at	#N/A	OslMA2	MAIAKSECERLAWALLLESNLVGNRNSNGDDDDDVDAAPAA
Os.27982.1.S1_at	#N/A	Unknown function	MOKDAPSPPLPPPVRYCGLEDYEQVDEEPPSPKSPSRKLSAAYRAELRRONALININGPRYRFRPGEKRRKKKKHNKSKHHTPPPSPTPRSPSPSPICPYEEYPTFEPDDPVMMRQSTMAYAEVLEHYNAADVGGGGVKYELVRAIFSGAIFTCKAAYGHLFFAEVRKDCKRYIPTCLWSDLDEADRVGGAGADPQVLDPEITSPSRRNYWFSCCDDEMKHPIKDGTSYHAGHL
Os.51491.1.S1_at	#N/A	Unknown function	MDISRTNEGYTSCGPVVEMSWHRAGVLLGLAQWPWLPGVSGILAVARVGFLVLSMARMGWELCHVFCPYAVRSVNAIKIQPPKNKTALQEMIGKINFRRFISNLSGRLEPTPLRLKADOKFTWGAEQQKALDSIKEYLSSPPVLIPOPQKLRYHLLSNECTVICKADVVKYMLASPISKGRVCKWIFSLETEFDLRYESPKAIGQQAIAFDIVDHDRNSIGSVEIVPWTLFFDGSVCTHGGIGLVIIARGASFEFAYTIKPYTTNQAEYEAVLKGQLLKEVEADVVEIMDDSLVISQLAGEYECKNDLVMYNEKCRELLRGFQLVISKHVSREQNVEANDLAQGASRYKPMKDVRVEATLSANDWRYDVHHYLNQPSQSASRKLRYKALRHTLDDLEYRTIDGFLKLCSADQTKVAIGDT
Os.5436.1.S1_at	#N/A	Unknown function	MPGSLPGGPVPHDKRSQDFSDNDVNLGLCERYILDYFLVVKRSSIDLVRERIDYPMFYPGRALKIKDRFALFMMMSLFLVSKYTDEVLMRVIETRSISYFAKLQRGDWSPHGTARKLKIRGPSPSEGIRSLSFLMEVSAWCRCVNCETC*